

DNA chimerism and its consequences after allogeneic hematopoietic cell transplantation

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The unphysiological formation of biological chimeras after allogeneic hematopoietic cell transplantation is not free of consequences. Recent findings suggest that in the transplant recipient some epithelial cells reveal, unexpectedly, donor-derived genotype and/or acquire genomic alterations. Since both phenomena are presented in the host epithelium, one could argue that they might be etiologically linked through a common background mechanism. We recently proposed that the incessant charge of the transplant recipient with donor-DNA and its integration in host epithelium by horizontal DNA transference may indeed be operative in the generation of epithelial cells with donor derived genome. On the other hand, the incessant incorporation of the foreign DNA into the host genome may result in genomic alterations. Lymphocyte-epithelial interactions between the two genetically distinct cell populations in the transplant recipient should be investigated more precisely not only in cellular but also in molecular level.

Allogeneic hematopoietic cell transplantation (allo-HCT) in humans results in true biological chimeras. While circulating hematopoietic cells and their tissue derivatives (e.g. Langerhans cells) become donor genotype after transplantation, other cells remain recipient in origin. This unphysiological formation of biological chimeras is not free of consequences. Recent findings of our group and others have shown that, besides Graft versus Host Disease (GvHD), there are also

other consequences in the co-existence of two genetically distinct populations in the transplant recipient. First, epithelial cells with donor-derived genotype emerge,^{1,2} a phenomenon, which was initially misinterpreted and falsely described as “stem cell plasticity”. Second, epithelial tissues of the host acquire genomic alterations.³

Is chimerism in epithelium after allo-HCT, namely the emergence of distinct epithelial cells containing donor-derived genome, a real phenomenon or solely a technical artefact? Despite the initial scepticism and the methodological limitations on the detection of donor-derived non-hematopoietic cells in the transplant recipient, more recent studies of our group and others using strict criteria and examinations of isolated single cells clearly confirmed that following allo-HCT in humans, epithelial cells with donor-derived genotype emerge.^{4,5} How does epithelial chimerism after allo-HCT occur? The mechanisms underlying this phenomenon remain unclear and divergent. Suggested mechanisms attempting to explain epithelial chimeric events after allo-HCT include transdifferentiation of hematopoietic cells into epithelial cells, generation of epithelial cells from unknown epithelial precursors and/or universal stem cells in the graft, and fusion of donor hematopoietic cells with recipient epithelial cells.⁶⁻⁹ More recent findings suggest molecule trafficking as a novel mechanism of epithelial chimerism after allo-HCT. Jang et al¹⁰ found that when murine hematopoietic stem cells are co-cultured with injured liver separated by a barrier, they may convert into liver-like

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cells, Aliotta et al¹¹ and Ratajczak J et al¹² showed that this phenotypical conversion may be due to mRNA transfer between cells, resulting in an aberrant expression of foreign proteins in the recipient hematopoietic cells. However, such an mRNA transfer from epithelial to hematopoietic cells cannot explain recent findings of our group and others after clinical transplantation. First, Y-chromosome positive epithelial-like cells found in female allotransplant recipients were negative for expression of hematopoietic markers. Second, large amount of donor-DNA has been detected in blood-free fingernails obtained from transplanted recipients.¹³ We also evaluated by quantitative microsatellite analysis the amount of donor DNA in 176 buccal swabs obtained from 71 patients after allogeneic transplantation and we found a high amount of donor-DNA (mean 26.6%) in the majority (89.7%) of them although no donor hematopoietic cells were evident in the samples by immunofluorescence.¹⁴

We recently proposed horizontal DNA transference as an alternative explanation for epithelial chimerism after allo-HCT.¹⁴ Production of donor cells from the engrafted bone marrow is an ongoing process in the allo-transplanted recipient. Apoptosis is a well-recognized source of DNA in several clinical settings, such as cancer, extensive burning, GvHD and transplantation.¹⁵⁻²⁰ Donor cells undergoing apoptosis release donor-DNA packaged into apoptotic bodies.²¹ Although foreign DNA is normally cleared up,²² the fate of the large amount of released donor-derived genetic material in the transplant recipient is unknown. In an in vitro co-culture system mimicking the lymphocyte-epithelial interaction we showed that DNA can be horizontally transferred from apoptotic hematopoietic cells to the cytoplasm and nucleus of epithelial cell lines through phagocytosis of apoptotic bodies.¹⁴ Both lysosomal inhibition in epithelial cells and repetitive load with apoptotic bodies, which may lead to saturation of lysosomal activity, increased the intercellular and intranuclear DNA delivery. The incessant charge of the transplant recipient with donor-DNA obtained from the engrafted bone marrow and its

illegitimate integration in host epithelium by horizontal gene transfer may indeed be operative in the generation of epithelial cells with donor derived genome in transplant recipients.

Horizontal gene transfer is well described in prokaryotic organisms as a mechanism for functional and phenotypic change in order to adapt to different environmental conditions. Horizontal gene transfer has also been shown in eukaryotic organisms both in vitro and in vivo. In vitro, the group of Holmgren has demonstrated that genomic DNA can be transferred to phagocytosing cells via the uptake of apoptotic cells.²³⁻²⁴ In vivo, large fragments of plasmid DNA fed to mice passed into the bloodstream and integrated into the nucleus of white blood cells, spleen and liver cells.^{25, 26} A critical role of DNA trafficking between cells has been also suggested in different clinical settings as in inflammation, in cancer and after organ transplantation.^{17, 18, 27, 28} Most recently, Ehnfors et al²⁹ used a rat-to-SCID mouse tumor xenotransplant system and showed that tumor DNA can be transferred in vivo by detecting rat and mouse fusion chromosomes in mouse tumor stromal cells. Such an approach could be also applicable in a hematopoietic cell xenotransplantation system (e.g. human-to-mouse) where the detection of human-mouse fusion chromosomes would show that horizontal gene transfer can be operational in the context of allo-HCT.

After allo-HCT, epithelial tissues become injured through the preparative regimen and are then potentially attacked by allo-reactive T cells. The net effect of these alloantigenic reactions is tissue stress and apoptosis, which we recognize clinically as GvHD. We hypothesized that chronic tissue stress due to interaction of donor-derived lymphocytes with host epithelium in the biological chimeras developed after HCT may cause genomic alterations. Indeed, our group and others found frequent genomic alterations measured as microsatellite instability (MSI, an established indicator of general genomic instability) in non-neoplastic epithelial tissues of patients who underwent allo-HCT, but not in control subjects including healthy individuals, autologous

transplanted patients and patients before allo-HCT.^{3, 30} Analysis of clinical data suggested the pivotal role of GvHD in the occurrence of MSI after allo HCT.³¹ In an in vitro mutation analysis system we found that allostimulated Mixed Lymphocyte Cultures may induce MSI in co-cultured epithelial cells.³¹

The development of epithelial cells with donor-derived genotype and the accumulation of genomic alterations in the epithelial tissues are only two of these recently recognized phenomena occurring in the chimeric organisms after allogeneic HCT. Since both phenomena are presented in the host epithelium, one could argue that they might be etiologically linked through a common background mechanism. A possible link between these two phenomena could be provided by horizontal gene transfer.

Taken together, current research results could support the following scenario after allo-HCT (**Figure 1**): the engrafted bone marrow produces continuously hematopoietic cells which after their programmed death charge constantly the host environment with donor-derived apoptotic bodies. In this context, the excessive amount of foreign material taken up repetitively by the recipient's professional and non-professional phagocytic cells may overwhelm their lysosomal capacity and thus part of the donor-derived apoptotic DNA fragments may escape degradation in the cytoplasm, be transferred into the nucleus and integrate within the recipient genome (DNA chimerism). This phenomenon may be more pronounced in GvHD lesions where the generated reactive oxygen species (ROS) from the activated lymphocytes may destabilize and damage lysosomes. Furthermore, the incorporation of the foreign DNA into the host genome could result in physical rearrangements at the site of integration, including point mutations, deletions, interruptions of coding sequences and chromosomal breakages. This "inappropriate" illegitimate integration of donor DNA in epithelial cells after allogeneic HCT may come in light as detection of epithelial cells with donor-derived genotype or as genomic instability in the epithelium and may have implications in the development of secondary cancers.

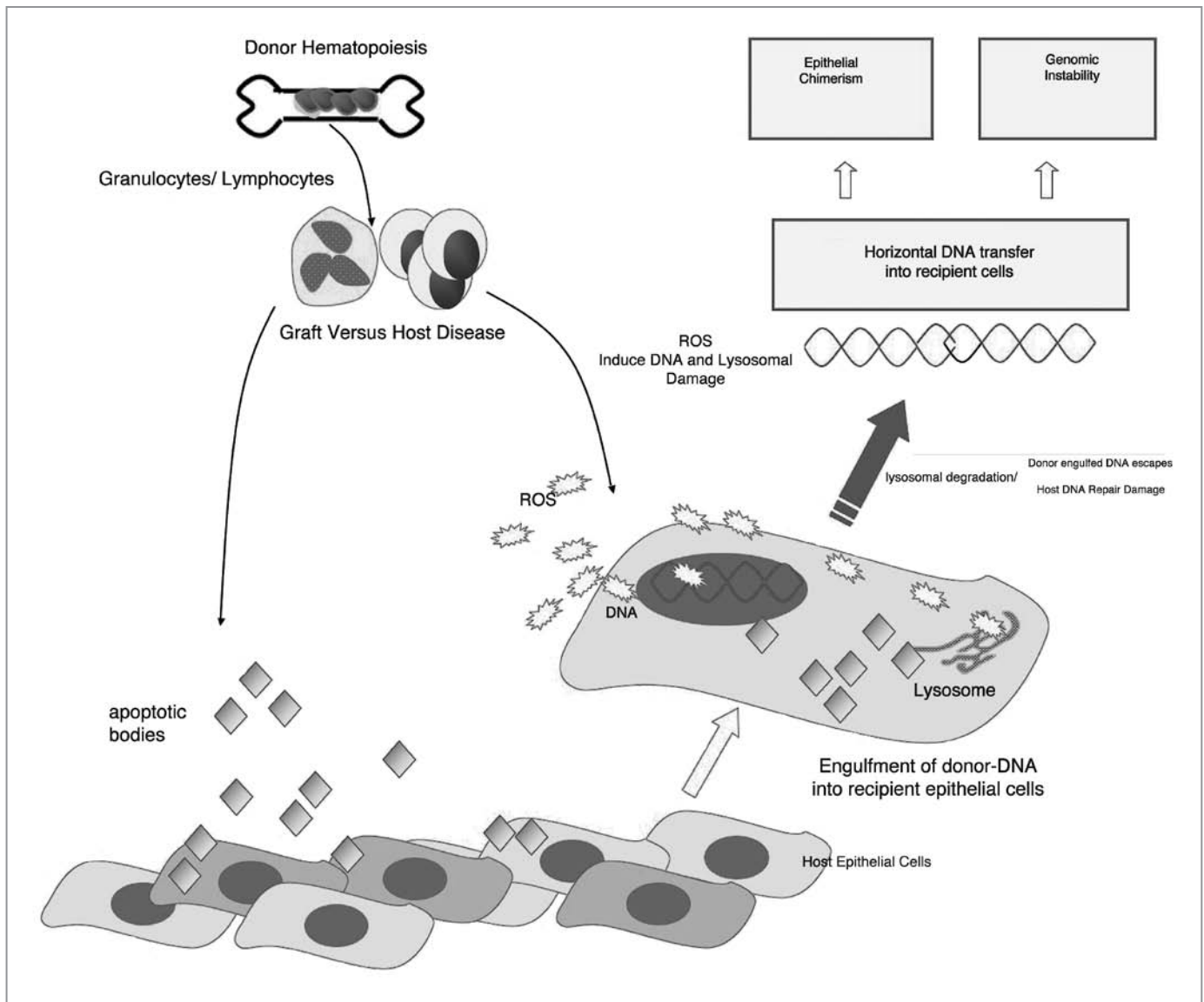


Figure 1. A proposed model of DNA Chimerism through Horizontal Gene Transfer and its consequences after allogeneic hematopoietic cell transplantation (see text).

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